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# EVIDENCE FOR THE METABOLIC FORMATION OF A VICINAL DIHYDRODIOL-EPOXIDE FROM THE POTENT MUTAGEN 1-NITROBENZO(A)PYRENE

MING W. CHOU AND PETER P. FU

National Center for Toxicological Research, Jefferson, AR 72079

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Summary: Metabolism of 1-nitrobenzo(a)pyrene (1-nitro-BaP) by rat liver microsomes yielded 1-nitro-BaP trans-7,8-dihydrodiol, 1-nitro-BaP trans-9,10-dihydrodiol and 1-nitro-BaP 7,8,9,10-tetrahydrotetrol. Formation of these metabolites suggests that a vicinal 7,8,9,10-dihydrodiol-epoxide is a metabolite of 1-nitro-BaP.

Nitro-polycyclic aromatic hydrocarbons (nitro-PAHs) have recently been identified as a new class of potent mutagenic and carcinogenic environmental pollutants existing in fly ash, diesel emissions, photocopier toners, cigarette smoke and soils (1-5). Since many nitro-PAHs are widespread in the environment and are carcinogenic in experimental animals (5,6) a major concern now is the possible hazard of these compounds to human health. Published results indicated that both reduction of the nitro substituent of nitro-PAHs (7,8) and ring oxidation (9-12) can be involved in the metabolic activation. However, it is not known if nitro-PAHs can be metabolized to vicinal diolepoxides, the activated form of PAHs. 1-Nitro-BaP, together with the 3- and 6-nitro-BaP isomers, has been found to be formed in model atmospheres containing trace quantities of BaP, nitrogen oxide and nitric acid (13). report that the in vitro aerobic metabolism of 1-nitro-BaP, a potent bacterial mutagen both in the presence and absence of mammalian liver homogenate activation enzymes (14), yields trans-7.8- and 9.10-dihydrodiols and 7.8.9.10tetrahydrotetrol. These results suggest that a vicinal dihydrodiol-epoxide can be a metabolite of a nitro-PAH.

<sup>&</sup>lt;u>Abbreviations</u>: Nitro-PAH, nitro-polycyclic aromatic hydrocarbon; BaP, benzo(a)-pyrene; 1-nitro-BaP trans-7,8-dihydrodiol, trans-7,8-dihydroxy-1-nitro-7,8-dihydro-BaP; 1-nitro-BaP 7,8,9,10-tetrahydrotetrol, 7,8,9,10-tetrahydroxy-1-nitro-7,8,9,10-tetrahydro-BaP; HPLC, high performance liquid chromatography; NMR, nuclear magnetic resonance.

## MATERIALS AND METHODS

Synthesis of 1-Nitrobenzo(a)pyrene: 9,10-Dihydrobenzo(a)pyren-7(8H)-one (Aldrich Chem. Co.) was reduced to 7,8,9,10-tetrahydro-BaP via Wolff-Kishner reduction (15). Nitration of 7,8,9,10-tetrahydro-BaP with sodium nitrate in trifluoroacetic acid and trifluoroacetic anhydride at ambient temperature under argon gave a mixture of 1-, 3- and 6-nitro-7,8,9,10-tetrahydro-BaP in good yield (16). The reaction mixture was separated by HPLC employing a DuPont Zorbax SIL column (9.4 x 250 mm) and eluting with 2% tetrahydrofuran in hexane in a flow rate of 3 ml/min. The retention time of 1-nitro-7,8,9,10-tetrahydro-BaP was 18.2 min. Mass spectral analysis indicated molecular ion at m/z 301. Dehydrogenation of 1-nitro-7,8,9,10-tetrahydro-BaP with dichlorodicyanobenzo-quinone (17) gave 1-nitro-BaP which was purified by chromatography on a Florisil column with benzene as the eluting solvent; mp 250-250.5°, m/z of M at 297; 500 MHz proton NMR (acetone-d<sub>6</sub>)  $\delta$  7.94 (dd,1,H-8), 8.02 (dd,1,H-9), 8.19 (d,1,H-4), 8.36 (d,1,H-3), 8.39 (d,1,H-5), 8.49 (d,1,H-7), 8.78 (d,1,H-2), 8.94 (s,1,H-6), 9.12 (d,1,H-12), 9.32 (d,1,H-10), 9.57 (d,1,H-11); J<sub>2</sub>, 3 = 8.3, J<sub>4,5</sub> = 9.2, J<sub>11,12</sub> = 9.6 Hz.

In Vitro Incubations: Male Sprague-Dawley rats (80-100 g), obtained from our breeding colony, received intraperitoneal injections of 3-methylcholanthrene (25 mg/kg body weight) on 3 consecutive days before sacrifice. Liver microsomes were prepared as previously described (18). Incubation mixtures contained 50 mmol Tris-HCl, pH 7.5, 3 mmol MgCl<sub>2</sub>, 1 mmol NADP<sup>†</sup>, 2 mmol glucose-6-phosphate, 100 units glucose-6-phosphate dehydrogenase, 1 g microsomal protein and 40  $\mu$ mol 1-nitro-BaP (dissolved in 40 ml acetone) in a total incubation volume of 1 l. Incubations were conducted aerobically with shaking for 60 min at 37° and then quenched by the addition of 1 l acetone. The metabolites and residual substrate were partitioned into 2 l ethyl acetate, and the organic phase was dried with anhydrous sodium sulfate. The solvent was evaporated under reduced pressure and the residue was dissolved in 1 ml methanol for analysis by HPLC.

Separation and Characterization of Metabolites: Reversed-phase HPLC was performed with a Beckman system consisting of two model 100A pumps, a model 210 injector, a model 420 solvent programmer and a Waters Associates model 440 absorbance (254 nm) detector. Metabolites were separated by using a DuPont Zorbax ODS column (9.4 mm x 25 cm) and eluting with a 20-min linear gradient of 50-100% methanol in water at a flow rate of 3 ml/min.

Uv-visible spectra were obtained with a Beckman model 25 spectrometer. Mass spectra were recorded with a Finnigan model 4000 system. The samples were introduced with a solid probe and ionized at 70 eV with an ionizer temperature of  $250^{\circ}$ . <sup>1</sup>H NMR spectra were obtained with a Bruker WM 500 spectrometer.

#### RESULTS AND DISCUSSION

The HPLC profile of the ethyl acetate extractable metabolites obtained from incubation of 1-nitro-BaP with rat liver microsomes is shown in Figure 1. The chromatographic peak  $\underline{4}$  contained the recovered substrate 1-nitro-BaP. The uv-visible absorption spectra of the metabolites contained in chromatographic peaks  $\underline{2}$  and  $\underline{3}$  are shown in Fig. 2A. Both of these metabolites had similar mass spectra (Fig. 3) with a molecular ion at m/z 331 and a major fragment at m/z 267 due to loss of a water molecule and NO $_2$ . The structures of these metabolites were determined by analysis of their high resolution 500 MHz proton NMR

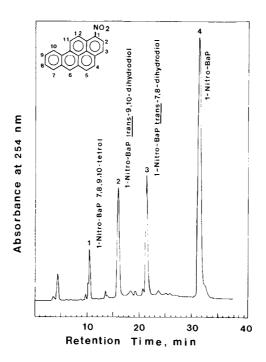


Fig. 1. Reversed-phase HPLC profile of ethyl acetate extractable metabolites obtained from incubation of 1-nitro-BaP with liver microsomes from 3-methylcholanthrene-treated male Sprague-Dawley rats. The chromatographic conditions are described in Materials and Methods.

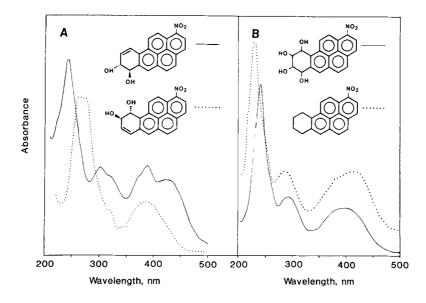


Fig. 2. Ultraviolet-visible absorption spectra (methanol) of (A) the metabolites identified as 1-nitro-BaP trans-7,8-dihydrodiol (contained in peak 3 of Fig. 1) (-----) and 1-nitro-BaP-9,10-dihydrodiol (contained in peak 2 of Fig. 1) (------) and (B) the metabolite contained in peak 1 of Fig. 1 identified as 1-nitro-BaP 7,8,8,10-tetrahydrotetrol (-----) and the synthetic compound 7,8,9,10-tetrahydro-1-nitro-BaP (------).

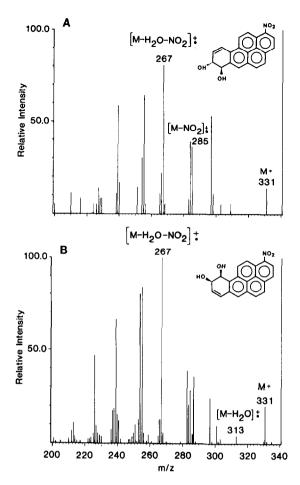


Fig. 3. Mass spectra of the metabolites identified as (A) 1-nitro-BaP trans-7,8,-dihydrodiol and (B) 1-nitro-BaP trans-9,10-dihydrodiol.

spectra (Fig. 4). The NMR resonance assignments were determined both by comparison to BaP  $\underline{\text{trans}}$ -7,8- and 9,10-dihydrodiols (19) and by extensive homonuclear decoupling experiments. Based on the mass and NMR spectral analysis, the metabolites contained in peaks  $\underline{2}$  and  $\underline{3}$  were identified as 1-nitro-BaP  $\underline{\text{trans}}$ -9,10-dihydrodiol and 1-nitro-BaP  $\underline{\text{trans}}$ -7,8-dihydrodiol, respectively. The proton NMR assignmens are as follows: 1-Nitro-BaP  $\underline{\text{trans}}$ -7,8-dihydrodiol (acetone-d<sub>6</sub> with trace D<sub>2</sub>0): 4.67 (dt,1, H<sub>8</sub>), 5.16 (d,1,H<sub>7</sub>), 6.44 (dd,1,H<sub>9</sub>), 7.65 (dd, 1,H<sub>10</sub>), 8.32 (d,1,H<sub>4</sub>), 8.44 (m, 2,H<sub>3,5</sub>), 8.71 (d,1,H<sub>2</sub>), 8.74 (s,1, H<sub>6</sub>) and 8.85 ppm (AB,2,H<sub>11,12</sub>); J<sub>2,3</sub> = J<sub>4,5</sub> = 8.6, J<sub>7,8</sub> = 11.2, J<sub>8,9</sub> = J<sub>8,10</sub> = 2.2 and J<sub>9,10</sub> = 10.3 Hz. 1-Nitro-BaP  $\underline{\text{trans}}$ -9,10-dihydrodiol (acetone-d<sub>6</sub> with trace D<sub>2</sub>0): 4.58 (dd,1,H<sub>9</sub>), 5.79 (d,1,H<sub>10</sub>), 6.46 (dd,1,H<sub>8</sub>), 7.03 (d,1,

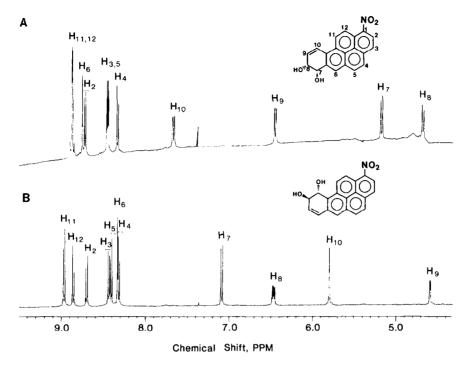


Fig. 4. 500 MHz proton NMR spectra of metabolites identified as (A) 1-nitro-BaP trans-7,8-dihydrodiol and (B) 1-nitro-BaP trans-7,8-dihydrodiol and (B) 1-nitro-BaP-trans-9,10-dihydrodiol. Chemical shifts are in ppm relative to tetramethylsilane.

 $H_7$ ), 8.31 (d,1, $H_4$ ), 8.33 (s,1, $H_6$ ), 8.41 (d,1, $H_5$ ), 8.43 (d,1, $H_3$ ), 8.70 (d,1,  $H_2$ ), 8.86 (d,1, $H_{12}$ ) and 8,96 ppm (d,1, $H_{11}$ );  $J_{2,3}$  = 8.6,  $J_{4,5}$  = 9.0,  $J_{7,8}$  = 9.5,  $J_{8,9}$  = 5.6,  $J_{9,10}$  = 2.2 and  $J_{11,12}$  = 9.9 Hz.

For 1-nitro-BaP trans-7,8-dihydrodiol, the coupling constants between the carbinol protons ( $J_{7,8}=11.2~\text{Hz}$ ) and between the non-benzylic olefinic and carbinol protons ( $J_{8,9}=2.2~\text{Hz}$ ) clearly indicate that this trans-dihydrodiol preferentially adonts a quasidiequatorial conformation (20). However, for 1-nitro-BaP trans-9,10-dihydrodiol, the coupling constants between the carbinol protons ( $J_{9,10}=2.2~\text{Hz}$ ) and between the non-benzylic olefinic and carbinol protons ( $J_{8,9}=5.6~\text{Hz}$ ) indicate that this trans-dihydrodiol preferentially adopts a quasidiaxial conformation (21). Thus, the preference of the dihydrodiol conformation for the 1-nitro-BaP is the same as that of the respective BaP dihydrodiols (20,21).

The material contained in peak  $\underline{1}$  had a uv-visible spectrum (Fig. 2B) that was similar to that of 7,8,9,10-tetrahydro-1-nitro-BaP, a synthetic standard.

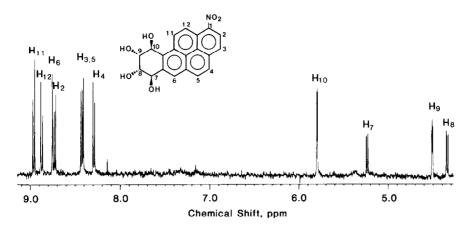


Fig. 5. 500 MHz proton NMR spectrum of metabolite identified as 1-nitro-BaP 7,8,9,10-tetrahydrotetrol.

This metabolite had a mass spectrum with molecular ions at m/z 365 and fragment ions at m/z 335 (loss of NO), 301 (loss of NO<sub>2</sub> and H<sub>2</sub>O) and 299 (loss of NO and 2H<sub>2</sub>O). These data were consistent with the metabolite being a 1-nitro-Bap 7.8.9.10-tetrahydrotetrol. A high resolution 500 MHz proton NMR spectrum of this metabolite confirmed the structural assignment (Fig. 5). The proton NMR assignments are as follows: (acetone- $d_6$  with trace  $D_2$ 0): 4.34 (dd,1, $H_8$ ), 4.51 (apparent  $t,1,H_0$ ), 5.24  $(d,1,H_7)$ , 5.80  $(d,1,H_{10})$ , 8.29  $(d,1,H_4)$ , 8.42  $(m,2,H_3,5)$ , 8.73  $(d,1,H_2)$ , 8.75  $(s,1,H_6)$ , 8.87  $(d,1,H_{12})$  and 8.96 ppm  $(d,1,H_1,H_2)$  $H_{11}$ );  $J_{2,3} = 8.6$ ,  $J_{4,5} = 9.0$ ,  $J_{7,8} = 9.0$ ,  $J_{8,9} = 2.2$ ,  $J_{9,10} = 3.4$  and  $J_{11,12} = 9.0$ 9.9 Hz. The observed coupling constants of  $J_{7.8}$ ,  $J_{8.9}$  and  $J_{9.10}$  indicate that this tetrol preferentially adopts a half chair conformation and has a transcis-trans configuration between  $H_7-H_8$ ,  $H_8-H_9$  and  $H_9-H_{10}$ , respectively. avoid steric interaction with the bay region hydrogen at  $C_{11}$  (22), the hydroxyl substituent at  $C_{10}$  maintains a quasidiaxial conformation. Consequently, the hydroxyl groups at  $C_7$ ,  $C_8$  and  $C_9$  are at quasiequatorial, quasiequatorial and quasiaxial, respectively.

Although <u>trans</u>-dihydrodiols have been identified as metabolites of 1-nitropyrene (11), 6-nitrochrysene (23) and 7-nitrobenz[a]anthracene (24), it was not known if a vicinal dihydrodiol-epoxide could be enzymatically formed from metabolism of nitro-PAHs. In this report, we found 1-nitro-BaP is metabolized to a 7,8,9,10-tetrahydrotetrol which indicates that a vicinal dihydro-

diol-epoxide is a metabolite of 1-nitro-BaP. The trans-cis-trans configuration at the terminal benzo ring of the 1-nitro-BaP 7.8.9.10-tetrahydrotetrol metabolite implies that the metabolically formed vicinal dihydrodiol-epoxide is either 1-nitro-BaP trans-7,8-dihydrodiol anti-9,10-epoxide or 1-nitro-BaP trans-9,10-dihydrodiol anti-7,8-epoxide.

1-Nitro-BaP has been found to be a potent bacterial mutagen in Salmonella typhimurium in the presence of mammalian liver homogenate activation enzymes In the tester strains TA98 and TA100, the mutagenic activation of 1-nitro-BaP is five-fold higher than that of BaP which itself is also a potent bacterial mutagen (13). These results together with our findings reported in this paper suggest that the enzymatically formed vicinal dihydrodiol-epoxide may be an ultimate mutagenic metabolite of 1-nitro-BaP when tested in the S. typhimurium with activation enzymes. On the other hand, 1-nitro-BaP is also a direct-acting bacterial mutagen while BaP is not (14). The direct-acting mutagenic activity of 1-nitro-BaP may be due to the nitro reduction of this compound to the activated form, N-hyroxy-1-amino-BaP by the nitroreductase(s) present in the bacteria (5). Thus, there appear to be at least two unique metabolic activation pathways for 1-nitro-BaP as far as its mutagenicity is concerned. The first is nitro-reduction to an N-hydroxy arylamine which is the activated metabolite of aromatic amines. The second is the ring oxidation to a vicinal dihydrodiol-epoxide which is a common ultimate mutagenic and carcinogenic metabolite of PAHs.

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